```
           MGAGGRMQVSPPSKKSETDTIKRVPCETPPFTVGBLKKAIPPHCFKRSIPRSFSYLIWDIIIASCFYYVATTYFPLLPHPLSYFANPLYWACQGCVLTGV 100
                   10        20        30        40        50        60        70        80        90       100

SEQIDNO12  MGAGGRMQVSPPSKKSETDTIKRVPCETPPFTVGBLKKAIPPHCFKRSIPRSFSYLIWDIIIASCFYYVATTYFPLLPHPLSYFANPLYWACQGCVLTGV 100
SEQIDNO16  MGAGGRMQVSPPSKKSETDNIKRVPCETPPFTVGBLKKAIPPHCFKRSIPRSFSYLIWDIIIASCFYYVATTYFPLLPHPLSYFANPLYWACQGCVLTGV 100


           WVIAHECGHHAFSDYQWLDDTVGLIFHSFLLVXYFSWKYSHRRHHSNTGSLERDEVFVPKKKKSDIKWYGKYLNNPLGRTVMLTVQFTLGWPLYLAFNVSG 200
                  110       120       130       140       150       160       170       180       190       200

SBQIDNO12  WVIAHKCGHHAFSDYQWLDDTVGLIFHSFLLVXYFSWKYSHRRHHSNTGSLERDEVFVPKKKKSDIKWYGKYLNNPLGRTVMLTVQFTLGWPLYLAFNVSG 200
SBQIDNO16  WVIAHECGHHAFSDYQWLDDTVGLIFHSFLLVPYFSWKYSHRRHHSNTGSLERDEVFVPKKKKSDIKWYGKYHNNPLGRTVMLTVQFTLGWPLYLAFNVSG 200


           RPYDGGFACHFHPNAPIYNDRERLQIYISDAGILAVCYGLFRYAAXQGVASMVCFYGVPLLXVNGFLVLITYLQHTHPSLPHYDSSEWDWLRGALATVDR 300
                  210       220       230       240       250       260       270       280       290       300

SEQIDNO12  RPYDGGFACHFHPNAPIYNDRERLQIYISDAGILAVCYGLFRYAAXQGVASMVCFYGVPLLXVNGFLVLITYLQHTHPSLPHYDSSEWDWLRGALATVDR 300
SEQIDNO16  RPYDGGFACHFHPNAPIYNDRERLQIYISDAGILAVCYGLFRYAAAQGVASMVCFYGVPLLIVNGFLVLITYLQHTHPSLPHYDSSEWDWLRGALATVDR 300


           DYGILNKVFHNITDTHVAHHLFSTMPHYHAMEATKAIKPILGBYYQFDGTPVVKAMWREAKBCIYVEPDRQGEKKGVFWYNNKL 384
                  310       320       330       340       350       360       370       380

SEQIDNO12  DYGILNKVFHNITDTHVAHHLFSTMPHYHAMEATKAIKPILGBYYQFDGTPVVKAMWREAKBCIYVEPDRQGEKKGVFWYNNKL 384
SEQIDNO16  DYGILNKVFHNITDTHVAHHLFSTMPHYHAMEATKAIKPILGBYYQFDGTPVVKAMWREAKBCIYVEPDRQGEKKGVFWYNNKL 384
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